

Amendments to the Claims:

This Listing of Claims should replace all prior versions and listings of claims in this application.

Listing of Claims:

Claim 1 (Currently Amended): A method for detecting the presence or absence of a bacterium in a sample selected from a wound, a body fluid or fluid from a wound, said method comprising the steps of:

a) contacting a said sample with a detectably labeled synthetic serpin α 1-proteinase inhibitor reactive site loop domain peptide substrate under conditions that result in modification cleavage of said substrate by an enzyme produced in said sample by a bacterium; and

b) detecting a modification cleavage or an absence of the modification cleavage of the substrate, the modification cleavage of the substrate indicating the presence of the bacterium in the sample and absence of the modification cleavage of the substrate indicating absence of the bacterium in the sample.

Claim 2 (Original): A method according to Claim 1, wherein the bacterium is a wound-specific bacterium selected from the group consisting of *Staphylococcus aureus*, *Staphylococcus epidermidis*, *Streptococcus pyogenes*, *Pseudomonas aeruginosa*, *Enterococcus faecalis*, *Serratia marcescens*, *Proteus mirabilis*, *Enterobacter cloacae*, *Acetobacter anitratus*, *Klebsiella pneumonia*, and *Escherichia coli*.

Claim 3 (Currently Amended): A method according to Claims 1, wherein the enzyme is a protease.

Claim 4 (Previously Presented): A method according to Claim 1, wherein the substrate is labeled with a fluorescent probe and a quencher dye molecule.

Claim 5 (Previously Presented): A method according to Claim 1, wherein the substrate is labeled by a label selected from the group consisting of spin labels, antigen tags, epitope tags, haptens, enzyme labels, prosthetic groups, fluorescent materials, pH-sensitive materials, chemiluminescent materials, colorimetric components, bioluminescent materials, and radioactive materials.

Claim 6 (Previously Presented): A method according to Claim 5, wherein the substrate comprises at least one of the peptides selected from the group consisting of EAAGAMFLEAIPK (SEQ ID NO: 1), EGAMFLEAIPMSIPK (SEQ ID NO: 2), KGTEAAGAMFLEAIPMSIPPEVK (SEQ ID NO: 3), GAMFLEAIPMSIPPE (SEQ ID NO: 4), and CGAMFLEAIPMSIPAAHHHHH (SEQ ID NO: 5).

Claim 7 (Currently Amended): A method according to Claim 1, wherein the sample is selected from the group consisting of a wound surface on a subject and a body fluid from a wound on a subject.

Claim 8 (Previously Presented): A method according to Claim 1, wherein the substrate is on a solid support.

Claim 9. (Previously Presented): A method according to Claim 8, wherein the solid support is selected from the group consisting of a wound dressing, a container for holding body fluids, a disk, a scope, a filter, a lens, a foam, a cloth, a paper, a suture, a dipstick, a swab, a urine collection bag, a blood collection bag, a plasma collection bag, a test tube, a catheter, and a well of a microplate.

Claim 10 (Previously Presented): A method according to Claim 8, wherein the solid support comprises a material required to be free of microbial contaminants.

Claim 11 (Previously Presented): A method according to Claim 1, wherein the substrate comprises at least two dissimilar colorimetric components and the substrate is attached to a solid support, wherein modification of the substrate comprises cleaving at least a portion of the substrate that includes one of the colorimetric components, the cleaving resulting in a visible color change.

Claim 12 (Previously Presented): A method according to Claim 11, wherein the colorimetric components are covalently attached to the peptide.

Claims 13-22 (Canceled).